

RAW SEQUENCE LISTING

1 <110> APPLICANT: Luiten, Rosalie

PATENT APPLICATION: US/09/766,889C

DATE: 06/03/2003 TIME: 10:11:47

Input Set : N:\Crf4\05222003\1766889B.raw
Output Set: N:\CRF4\06032003\1766889C.raw

```
Boon-Falleur, Thierry
              van der Bruggen, Pierre
              Stroobant, Vincent
              Demotte, Nathalie
              Schultz, Erwin
     7 <120> TITLE OF INVENTION: MAGE ANTIGENIC PEPTIDES WHICH BIND HLA-B35 AND HLA-B44
     8 <130> FILE REFERENCE: L00461/70104
C--> 9 <140> CURRENT APPLICATION NUMBER: US/09/766,889C
     10 <141> CURRENT FILING DATE: 2001-01-19
     11 <150> PRIOR APPLICATION NUMBER: US 60/177,242
     12 <151> PRIOR FILING DATE: 2000-01-20
     13 <150> PRIOR APPLICATION NUMBER: US 60/243,212
     14 <151> PRIOR FILING DATE: 2000-10-25
     15 <160> NUMBER OF SEO ID NOS: 59
     16 <170> SOFTWARE: PatentIn 3.1
     18 <210> SEQ ID NO: 1
     19 <211> LENGTH: 930
     20 <212> TYPE: DNA
     21 <213> ORGANISM: Homo sapiens
     22 <400> SEQUENCE: 1
    23
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    24
                                                                                      120
              gaggecetgg geetggtgtg tgtgeagget geeaceteet eeteetetee tetggteetg
    25
              ggcaccetgg aggaggtgcc cactgetggg teaacagate etceccagag teetcaggga
                                                                                      180
    26
                                                                                      240
              geotecgect tteccactae cateaactte actegacaga ggcaacccag tgagggttee
    27
              ageageegtq aagaggaggg gecaageace tettgtatee tggagteett gtteegagea
                                                                                      300
    28
                                                                                      360
              qtaatcacta agaaggtggc tgatttggtt ggttttctqc tcctcaaata tcgagccagg
    29
              gagccagtca caaaggcaga aatgctggag agtgtcatca aaaattacaa gcactgtttt
                                                                                      420
    30
              ectgagatet teggeaaage etetgagtee ttgeagetgg tetttggeat tgaegtgaag
                                                                                      480
    31
                                                                                      540
              gaaqcaqacc ccaccggcca ctcctatgtc cttgtcacct gcctaggtct ctcctatgat
    32
              ggcctgctgg gtgataatca gatcatgccc aagacaggct tcctgataat tgtcctggtc
                                                                                      600
    33
                                                                                      660
              atgattgcaa tggagggcgg ccatgctcct gaggaggaaa tctgggagga gctgagtgtg
    34
              atggaggtgt atgatgggag ggagcacagt gcctatgggg agcccaggaa gctgctcacc
                                                                                      720
    35
              caaqatttqq tqcaqqaaaa qtacctqqaq taccqqcaqq tqccqqacaq tqatcccqca
                                                                                      780
    36
                                                                                      840
              egetatgaqt teetgtqqqq teeaaqqqee eteqetgaaa eeaqetatgt gaaaqteett
    37
              gagtatgtga tcaaggtcag tgcaagagtt cgctttttct tcccatccct gcgtgaagca
                                                                                      900
                                                                                      930
    38
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    40 <210> SEO ID NO: 2
    41 <211> LENGTH: 309
     42 <212> TYPE: PRT
     43 <213> ORGANISM: Homo sapiens
    44 <400> SEOUENCE: 2
    45
              Met Ser Leu Glu Gln Arg Ser Leu His Cys Lys Pro Glu Glu Ala Leu
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95 <212> TYPE: DNA

96 <213> ORGANISM: Homo sapiens

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Input Set : N:\Crf4\05222003\I766889B.raw Output Set: N:\CRF4\06032003\I766889C.raw

```
1
                                            1.0
46
         Glu Ala Gln Glu Ala Leu Gly Leu Val Cys Val Gln Ala Ala Thr
47
                                        25
                                                            30
48
                    20
          Ser Ser Ser Pro Leu Val Leu Gly Thr Leu Glu Glu Val Pro Thr
49
50
                                     40
          Ala Gly Ser Thr Asp Pro Pro Gln Ser Pro Gln Gly Ala Ser Ala Phe
51
52
                                55
53
          Pro Thr Thr Ile Asn Phe Thr Arg Gln Arg Gln Pro Ser Glu Gly Ser
                             70
                                                75
54
         Ser Ser Arg Glu Glu Glu Gly Pro Ser Thr Ser Cys Ile Leu Glu Ser
55
56
                         8.5
                                             90
          Leu Phe Arg Ala Val Ile Thr Lys Lys Val Ala Asp Leu Val Gly Phe
57
58
                     100
                                         105
                                                            110
          Leu Leu Leu Lys Tyr Arg Ala Arg Glu Pro Val Thr Lys Ala Glu Met
59
                                    120
                                                        125
60
          Leu Glu Ser Val Ile Lys Asn Tyr Lys His Cys Phe Pro Glu Ile Phe
61
                                                    140
62
            130
                                 135
         Gly Lys Ala Ser Glu Ser Leu Gln Leu Val Phe Gly Ile Asp Val Lys
63
64
          145
                            150
                                                155
         Glu Ala Asp Pro Thr Gly His Ser Tyr Val Leu Val Thr Cys Leu Gly
65
                         165
                                           170
                                                                175
66
67
          Leu Ser Tyr Asp Gly Leu Leu Gly Asp Asn Gln Ile Met Pro Lys Thr
68
                     180
                                        185
                                                            190
         Gly Phe Leu Ile Ile Val Leu Val Met Ile Ala Met Glu Gly Gly His
69
                                    200
70
                195
         Ala Pro Glu Glu Glu Ile Trp Glu Glu Leu Ser Val Met Glu Val Tyr
71
72
            210
                                 215
                                                    220
         Asp Gly Arg Glu His Ser Ala Tyr Gly Glu Pro Arg Lys Leu Leu Thr
73
74
          225
                             230
                                                235
         Gln Asp Leu Val Gln Glu Lys Tyr Leu Glu Tyr Arg Gln Val Pro Asp
7.5
76
                         245
                                            250
                                                                255
          Ser Asp Pro Ala Arg Tyr Glu Phe Leu Trp Gly Pro Arg Ala Leu Ala
77
7.8
                     260
                                        265
         Glu Thr Ser Tyr Val Lys Val Leu Glu Tyr Val Ile Lys Val Ser Ala
79
80
                                     280
          Arg Val Arg Phe Phe Pro Ser Leu Arg Glu Ala Ala Leu Arg Glu
81
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                                 295
82
83
         Glu Glu Glu Gly Val
         305
84
86 <210> SEQ ID NO: 3
87 <211> LENGTH: 31
88 <212> TYPE: DNA
89 <213> ORGANISM: Homo sapiens
90 <400> SEQUENCE: 3
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93 <210> SEQ ID NO: 4
94 <211> LENGTH: 30
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30

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DATE: 06/03/2003 PATENT APPLICATION: US/09/766,889C TIME: 10:11:47

Input Set : N:\Crf4\05222003\I766889B.raw Output Set: N:\CRF4\06032003\I766889C.raw

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100 <210> SEO ID NO: 5
101 <211> LENGTH: 12
102 <212> TYPE: PRT
103 <213> ORGANISM: Homo sapiens
104 <400> SEQUENCE: 5
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106
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108 <210> SEQ ID NO: 6
109 <211> LENGTH: 10
110 <212> TYPE: PRT
111 <213> ORGANISM: Homo sapiens
112 <400> SEQUENCE: 6
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                          5
114
116 <210> SEQ ID NO: 7
117 <211> LENGTH: 9
118 <212> TYPE: PRT
119 <213> ORGANISM: Homo sapiens
120 <400> SEQUENCE: 7
121
         Asp Pro Thr Gly His Ser Tyr Val Leu
122
                          5
124 <210> SEQ ID NO: 8
125 <211> LENGTH: 9
126 <212> TYPE: PRT
127 <213> ORGANISM: Homo sapiens
128 <400> SEOUENCE: 8
129
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130
          1
132 <210> SEQ ID NO: 9
133 <211> LENGTH: 10
134 <212> TYPE: PRT
135 <213> ORGANISM: Homo sapiens
136 <400> SEOUENCE: 9
137
      Lys Glu Ala Asp Pro Thr Gly His Ser Tyr
138
140 <210> SEO ID NO: 10
141 <211> LENGTH: 8
142 <212> TYPE: PRT
143 <213> ORGANISM: Homo sapiens
144 <400> SEQUENCE: 10
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146
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148 <210> SEQ ID NO: 11
149 <211> LENGTH: 72
150 <212> TYPE: DNA
151 <213> ORGANISM: Homo sapiens
152 <400> SEQUENCE: 11
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- RAW SEQUENCE LISTING

DATE: 06/03/2003

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Input Set: N:\Crf4\05222003\1766889B.raw
Output Set: N:\CRF4\06032003\1766889C.raw

```
60
153
          atgtctgagt ccttgcagct ggtctttggc attgacgtga aggaagcaga ccccaccggc
                                                                                  72
154
          cactectatt ga
156 <210> SEQ ID NO: 12
157 <211> LENGTH: 23
158 <212> TYPE: PRT
159 <213> ORGANISM: Homo sapiens
160 <400> SEQUENCE: 12
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161
162
163
          Asp Pro Thr Gly His Ser Tyr
164
                      20
166 <210> SEO ID NO: 13
167 <211> LENGTH: 33
168 <212> TYPE: DNA
169 <213> ORGANISM: Homo sapiens
170 <400> SEQUENCE: 13
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173 <210> SEQ ID NO: 14
174 <211> LENGTH: 10
175 <212> TYPE: PRT
176 <213> ORGANISM: Homo sapiens
177 <400> SEQUENCE: 14
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178
179
                           -5
181 <210> SEQ ID NO: 15
182 <211> LENGTH: 30
183 <212> TYPE: DNA
184 <213> ORGANISM: Homo sapiens
185 <400> SEQUENCE: 15
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186
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188 <210> SEO ID NO: 16
189 <211> LENGTH: 9
190 <212> TYPE: PRT
191 <213> ORGANISM: Homo sapiens
192 <400> SEQUENCE: 16
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194
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196 <210> SEO ID NO: 17
197 <211> LENGTH: 9
198 <212> TYPE: PRT
199 <213> ORGANISM: Homo sapiens
200 <400> SEQUENCE: 17
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201
202
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204 <210> SEQ ID NO: 18
205 <211> LENGTH: 9
206 <212> TYPE: PRT
207 <213> ORGANISM: Homo sapiens
208 <400> SEOUENCE: 18
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RAW SEQUENCE LISTING

DATE: 06/03/2003 PATENT APPLICATION: US/09/766,889C TIME: 10:11:47

Input Set : N:\Crf4\05222003\I766889B.raw Output Set: N:\CRF4\06032003\I766889C.raw

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210
212 <210> SEO ID NO: 19
213 <211> LENGTH: 9
214 <212> TYPE: PRT
215 <213> ORGANISM: Homo sapiens
216 <400> SEQUENCE: 19
217
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          1
218
220 <210> SEQ ID NO: 20
221 <211> LENGTH: 10
222 <212> TYPE: PRT
223 <213> ORGANISM: Homo sapiens
224 <400> SEQUENCE: 20
    Met Glu Val Asp Pro Ile Gly His Leu Tyr
225
226
                          5
228 <210> SEQ ID NO: 21
229 <211> LENGTH: 9
230 <212> TYPE: PRT
231 <213> ORGANISM: Homo sapiens
232 <400> SEQUENCE: 21
233
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234
          1 5
236 <210> SEQ ID NO: 22
237 <211> LENGTH: 8
238 <212> TYPE: PRT
239 <213> ORGANISM: Homo sapiens
240 <400> SEQUENCE: 22
241 Tyr Arg Pro Arg Pro Arg Arg Tyr
242
244 <210> SEQ ID NO: 23
245 <211> LENGTH: 10
246 <212> TYPE: PRT
247 <213> ORGANISM: Homo sapiens
248 <400> SEQUENCE: 23
249 Ser Pro Ser Ser Asn Arg Ile Arg Asn Thr
250
252 <210> SEQ ID NO: 24
253 <211> LENGTH: 9
254 <212> TYPE: PRT
255 <213> ORGANISM: Homo sapiens
256 <400> SEOUENCE: 24
257 Val Leu Pro Asp Val Phe Ile Arg Cys
258 1 5
260 <210> SEO ID NO: 25
261 <211> LENGTH: 10
262 <212> TYPE: PRT
263 <213> ORGANISM: Homo sapiens
264 <400> SEQUENCE: 25
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RAW SEQUENCE LISTING ERROR SUMMARY PATENT APPLICATION: US/09/766,889C

DATE: 06/03/2003 TIME: 10:11:48

Input Set : N:\Crf4\05222003\1766889B.raw
Output Set: N:\CRF4\06032003\1766889C.raw

Please Note:

.

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:53; Xaa Pos. 1

Invalid Line Length:

The rules require that a line not exceed 72 characters in length. This includes spaces.

Seq#:1; Line(s) 7,23,24,25,26,27,28,29,30,31,32,33,34,35,36,37,38Seq#:3; Line(s) 91

Seq#:4; Line(s) 98 Seq#:11; Line(s) 153,154 Seq#:13; Line(s) 171 Seq#:15; Line(s) 186 VERIFICATION SUMMARY

PATENT APPLICATION: US/09/766,889C

DATE: 06/03/2003 TIME: 10:11:48

Input Set : N:\Crf4\05222003\1766889B.raw
Output Set: N:\CRF4\06032003\1766889C.raw

L:9 M:270 C: Current Application Number differs, Wrong Format L:493 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:53 after pos.:0